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Fig. 1

SEQ ID NO: 1

GGCAGCCGCAGACCGTGGCTGAGCATGGAGCTGTCCCCCGCAGTCCT
CCGGAGATGCTGGAGGAGTCGGATTGCCGTACCCCTGGAGCTGAAG
TCAGCCCCAGCAAGAAGATGTGGATTAAGCTCGGTCTGCTGCGC
TACATGGTGAAGCAGTTGGAGAATGGGGAGATAAACATTGAGGAGCTG
AAGAAAAATCTGGAGTACACAGCTCTGCTGGAAGCCGTACATA
GATGAGACACGGCAAATCTTGACACGGAGGACGAGCTGCAGGAGCTG
CGGTCAAGATGCCGTGCCCTCGGAGGTGCGGGACTGGCTGGCCTCCACC
TTCACCCAGCAGGCCGGGCAAAGGCCGCCAGCAGAGGAGAACCCC
AAGTTCCGAAGCATTGTGCACGCTGTGCAGGCTGGATCTCGTGGAA
CGGATGTTCCGGAGAACATACACCTCTGTGGATCTCTGGTGTGTTGATGTC
GCGGTTCTCAACTGTCTCAAGAACCTGGATCTCTGGTGTGTTGATGTC
TTTCCTTGAACCAGGCAGCAGATGACCATGCCCTGAGGACCATTGTT
TTGAGTTGCTGACTCGGCATAACCTCATCAGCCGCTTCAAGATTCCC
ACTGTGTTTGATGAGTTCTGGATGCCTGGAGACAGGCTATGGG
AAGTACAAGAACCTTACCAACAGATCCACGCAGCCGATGTTACC
CAGACAGTCCATTGCTTCTGCTCCGCACAGGGATGGTGCAGTCCTG
TCGGAGATTGAGCTCCTGGCCATCATCTTGCTGCAGCTATCCATGAT
TATGAGCACACGGGCACTACCAACAGCTTCCACATCCAGACCAAGTCA
GAATGTGCCATCGTGTACAATGATCGTTAGTGGAGACAGGCTATGGG
ATCAGCTCTGTTCCGATTGATGCAGGATGATGAGATGAACATTTC
ATCAACCTCACCAAGGATGAGTTGAGAACACTCCGAGCCCTGGTCATT
GAGATGGTGTGGCCACAGACATGTCTGCCATTCCAGCAAGTGAAG
ACCATGAAGACAGCCTGCAACAGCTGGAGAGGATTGACAAGCCAAAG
GCCCTGTCTACTGCTCCATGCTGACATCAGCCACCCAACCAAG
CAGTGGTGGTCCACAGCCGTTGGACAGAGTGGCCTGCCCTTCTCCA
CTCTGTGACCGCACTCCACTCTAGTGGCACAGTCTCAGATAGGGTTC
ATCGACTTCATTGTGGAGCCCACATTCTCTGTGCTGACTGACGTGGCA
GAGAAGAGTGTTCAGCCCCTGGCGATGAGGACTCCAAGTCTAAAAAC
CAGCCCAGCTTCAGTGGCGCAGCCCTCTGGATGTGGAAGTGGGA
GACCCCAACCCTGATGTGGTCAGCTTCGTTCCACCTGGTCAAGCGC
ATTCAAGGAGAACAGCAGAAATGGAAGGAACGGGAGCAAGTGGCATT
ACCAACCAGATGTCCATTGACGAGCTGTCCCCCTGTGAAGAACAGGCC
CCCCCATCCCCCTGCCGAAGATGAACACAACCAGAACGGGATCTGGAT
TAGCCCTGGGCTGGCCCAGGTCTTCATTGAGTCCAAAGTGTGTTGATG
TCATCAGCACCATCCATCAGGACTGGCTCCCCATCTGCTCCAAGGG
GCGTGGTGTGGAAGAAACAAACCCACCTGAAGGCCAAATGCCAGAGAT
TTGGGGTTGGGAAAGGGCCCTCCCCACCTGACACCCACTGGGTGC
ACTTTAATGTTCCGGCAGCAAGACTGGGAACCTCAGGCTCCAGTGG
TCACTGTGCCCATCCCTCAGCCTCTGGATTCTTCTCATGGCCAGGTGG
CTGCCAGGGAGCGGGAGCTTCCCTGGAGGCTCCAGGGCCTGGGGA

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Fig. 1 (continued)

AGGGTCAGAGATGCCAGCCCCCTGGGACCTCCCCCATCCTTTGCCT
CCAAGTTCTAACAAATACATTGGGGTCCCTCAGCCCCCACCC
CAGATCTTAGCTGGCAGGTCTGGGTGCCCTTCCTCCCCTGGGAAG
GGCTGGAATAGGATAGAAAAGCTGGGGTTTCAGAGCCCTATGTGTGG
GGAGGGGAGTGGATTCTTCAGGGCATGGTACCTTCTAGGATCTGGG
AATGGGTGGAGAGGACATCCTCTCACCCAGAATTGCGGGAATT

Fig. 2

SEQ ID NO: 2

MELSPRSPPPEMLEESDCPSPLEKSAPSKKWIKLRSLLRYMVKQLEN
GEINIEELKKNLEYTASLLEAVYIDETRQILDTEDELQELRSDAVPSE
VRDWLASTFTQQARAKRRRAEEKPKFRSIVHAVQAGIFVERMFRRTYT
SVGPTYSTAVLNCLKNLDLWCDFVFSLNQAADDHALRTIVFELLTRHN
LISRFKIPTVFLMSFLDALETGYGKYKNPYHNQIHAADVQTVCFCFL
RTGMVHCLSEIELLAIIFAAAIHDYEHTGTTNSFHIQTKSECAIVYND
RSVLENHHISSVFRLMQDDEMNIFINLTKEFVELRALVIEMVLATDM
SCHFQQVKTMKTALQQLERIDKPKALSLLHAADISHPTKQWLVHSRW
TKALMEEFFRQGDKEAELGLPFSPLCDRTSTLVAQSQIGFIDFIVEPT
FSVLTDVAEKSVQPLADEDSKSKNQPSFQWRQPSLDVEVGDPNPDVVS
FRSTWVKRIQENKQWKERAASGITNQMSIDE LSPCEEAPPSPAED
HNQNGNLD

Fig. 3

SEQ ID NO: 3

5' - TGTGGAAGTGGGAGACCC - 3'

Fig. 4

SEQ ID NO: 4

5' - CTGAATGCGCTTGACCCAG - 3'

Fig. 5

SEQ ID NO: 5

5' - ACCCTGATGTGGTCAGTTCTTCCA - 3'